



PTO/SB/08a/b (07-05)

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Substitute for form 1449A/B/PTO INFORMATION DISCLOSURE STATEMENT BY APPLICANT <i>(Use as many sheets as necessary)</i>				Complete if Known	
				Application Number	10/509275
				Filing Date	May 25, 2005
				First Named Inventor	Benedikt Brors
				Art Unit	1653
				Examiner Name	Not Yet Assigned
Sheet	1	of	1	Attorney Docket Number	VOSS-P01-011

U.S. PATENT DOCUMENTS					
Examiner Initials*	Cite No. ¹	Document Number	Publication Date MM-DD-YYYY	Name of Patentee or Applicant of Cited Document	Pages, Columns, Lines, Where Relevant Passages or Relevant Figures Appear
		Number-Kind Code ² (if known)			

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/BLS	CE1	International Preliminary Examination Report for PCT/EP03/03291 issued 8/4/2005	

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PTO/SB/08a/b (08-03)

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Substitute for form 1449A/B/PTO				Complete If Known	
INFORMATION DISCLOSURE STATEMENT BY APPLICANT <i>(Use as many sheets as necessary)</i>				Application Number	10/509275
				Filing Date	September 27, 2004
				First Named Inventor	Benedikt Brors
				Art Unit	Not Yet Assigned
				Examiner Name	Not Yet Assigned
Sheet	1	of	2	Attorney Docket Number	VOSS-P01-011

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/BLS/	CA	Bustin, "Absolute Quantification of mRNA Using Real-time Reverse Transcription Polymerase Chain Reaction Assays," Journal of Molecular Endocrinology 25:169-193 (2000)			
	CB	Case-Green et al., "Analysing Genetic Information with DNA Arrays," Current Opinion in Chemical Biology 2:404-410 (1998)			
	CC	Chen et al., "Ratio-based Decisions and the Quantitative Analysis of cDNA Microarray Images," Journal of Biomedical Optics 2(4):364-374 (1997)			
	CD	Chudin et al., "Assessment of the Relationship Between Signal Intensities and Transcript Concentration for Affymetrix GeneChip® Arrays," Genome Biology 3(1):0005.1-0005.10 (2001)			
	CE	DeRisi et al., "Exploring the Metabolic and Genetic Control of Gene Expression on a Genomic Scale," Science 278:680-686 (1997)			
	CF	DeRisi et al., "Use of a cDNA Microarray to Analyse Gene Expression Patterns in Human Cancer," Nature Genetics 14:457-460 (1996)			
	CG	Duggan et al., "Expression Profiling Using cDNA Microarrays," Nature Genetics Supplement 21:10-14 (1999)			
	CH	Eickhoff et al., "Normalization of Array Hybridization Experiments in Differential Gene Expression Analysis," Nucleic Acids Research 27(22):i-iii (1999)			
	CI	Freeman et al., "Quantitative RT-PCR: Pitfalls and Potential," BioTechniques 26(1):112-125 (1999)			
	CJ	Golub et al., "Molecular Classification of Cancer: Class Discovery and Class Prediction by Gene Expression Monitoring," Science 286:531-537 (1999)			
	CK	Hegde et al., "A Concise Guide to cDNA Microarray Analysis," BioTechniques 29:548-562 (2000)			
	CL	Hughes et al., "Expression Profiling Using Microarrays Fabricated by an Ink-jet Oligonucleotide Synthesizer," Nature Biotechnology 19:342-347 (2001)			
/BLS/	CM	Hughes et al., "Functional Discovery via a Compendium of Expression Profiles," Cell 102:109-126 (2000)			

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/BLS/	CN	Iyer et al., "The Transcriptional Program in the Response of Human Fibroblasts to Serum," Science 283:83-87 (1999)	
	CO	Khan et al., "DNA Microarray Technology: The Anticipated Impact on the Study of Human Disease," Biochimica et Biophysica Acta 1423:M17-M28 (1999)	
	CP	Lee et al., "Importance of Replication in Microarray Gene Expression Studies: Statistical Methods and Evidence from Repetitive cDNA Hybridizations," PNAS 97(18):9834-9839 (2000)	
	CQ	Lipshutz et al., "High Density Synthetic Oligonucleotide Arrays," Nature Genetics Supplement 21:20-24 (1999)	
	CR	Lockhart et al., "Expression Monitoring by Hybridization to High-density Oligonucleotide Arrays," Nature Biotechnology 14:1675-1680 (1996)	
	CS	Lockhart et al., "Genomics, Gene Expression and DNA Arrays," Nature 405:827-836 (2000)	
	CT	Newton et al., "On Differential Variability of Expression Ratios: Improving Statistical Inference about Gene Expression Changes from Microarray Data," Journal of Computational Biology 8:37-52 (2001)	
	CU	Schadt et al., "Analyzing High-density Oligonucleotide Gene Expression Array Data," Journal of Cellular Biochemistry 80:192-202 (2000)	
	CV	Schena "Genome Analysis with Gene Expression Microarrays," BioEssays 18(5):427-431 (1996)	
	CW	Schena et al., "Quantitative Monitoring of Gene Expression Patterns with a Complementary DNA Microarray," Science 270:467-470 (1995)	
	CX	Schuchhardt et al., "Normalization Strategies for cDNA Microarrays," Nucleic Acids Research 28(10):i-v (2000)	
	CY	Shalon et al., "A DNA Microarray System for Analyzing Complex DNA Samples Using Two-color Fluorescent Probe Hybridization," Genome Research 6:639-649 (1996)	
	CZ	Spellman et al., "Comprehensive Identification of Cell Cycle-regulated Genes of the Yeast Saccharomyces cerevisiae by Microarray Hybridization," Molecular Biology of the Cell 9:3273-3297 (1998)	
	CA1	T. Beißbarth et al., "Processing and Quality Control of DNA Array Hybridization Data," Bioinformatics 16(11):1014-1022 (2000)	
	CB1	Velculescu et al., "Analysis of Human Transcriptomes," Nature Genetics 23:387-388 (1999)	
	CC1	Velculescu et al., "Serial Analysis of Gene Expression," Science 270:484-487 (1995)	
/BLS/	CD1	Yue et al., "An Evaluation of the Performance of cDNA Microarrays for Detecting Changes in Global mRNA Expression," Nucleic Acids Research 29(8):1-9 (2001)	

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